

Fig. 1

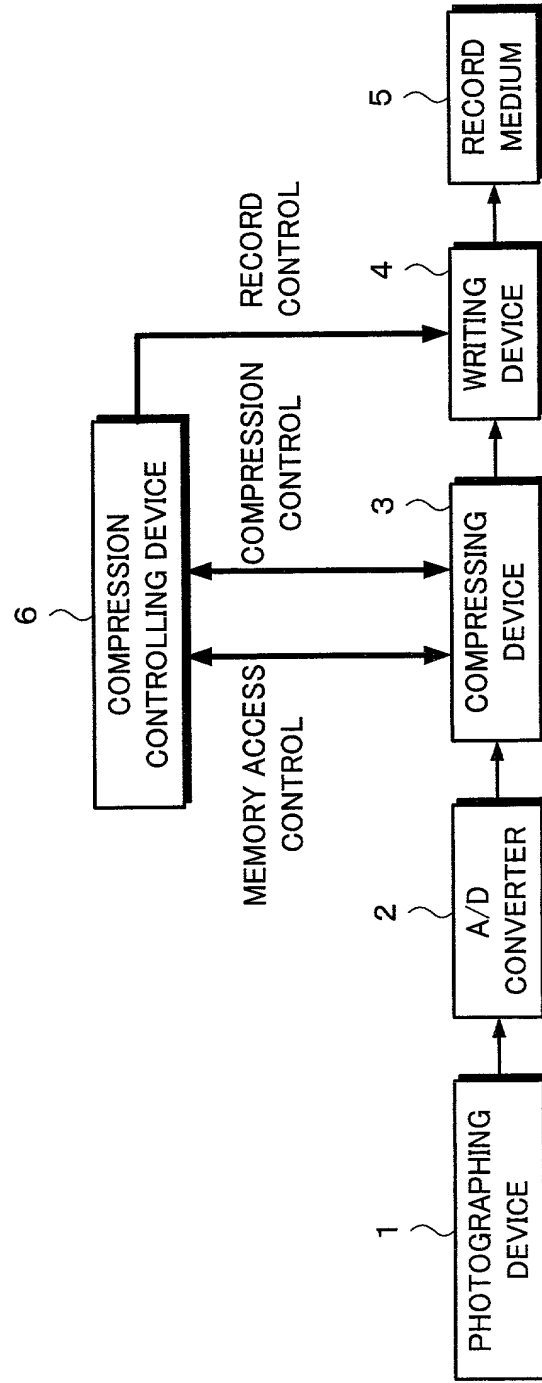


Fig. 3A

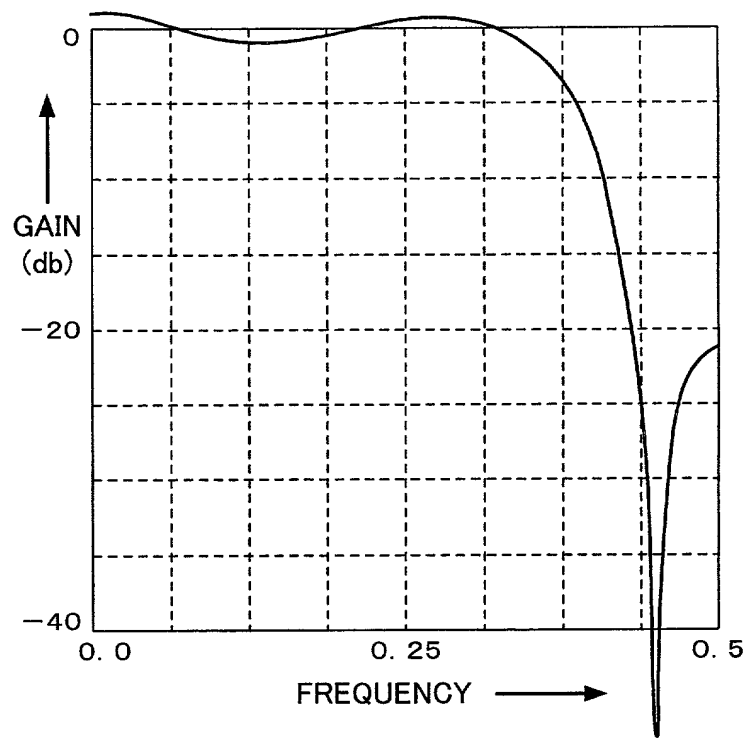


Fig. 3B

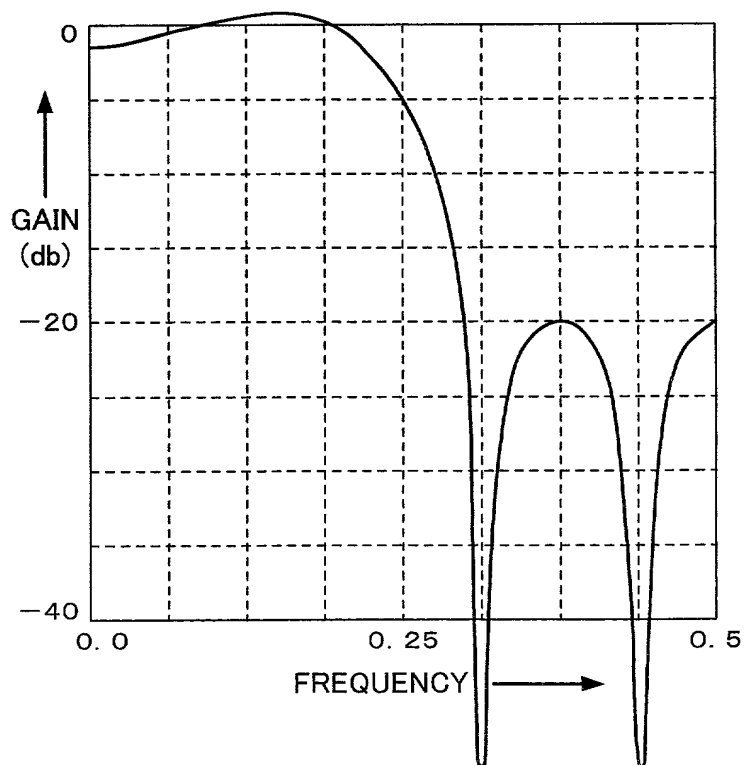


Fig. 4A

Fig. 4B

Fig. 4C

A diagram showing a rectangular frame with a width dimension of 352 PIXELS and a height dimension of 480 LINES. The dimensions are indicated by arrows pointing to the top and left edges of the frame.

Fig. 5

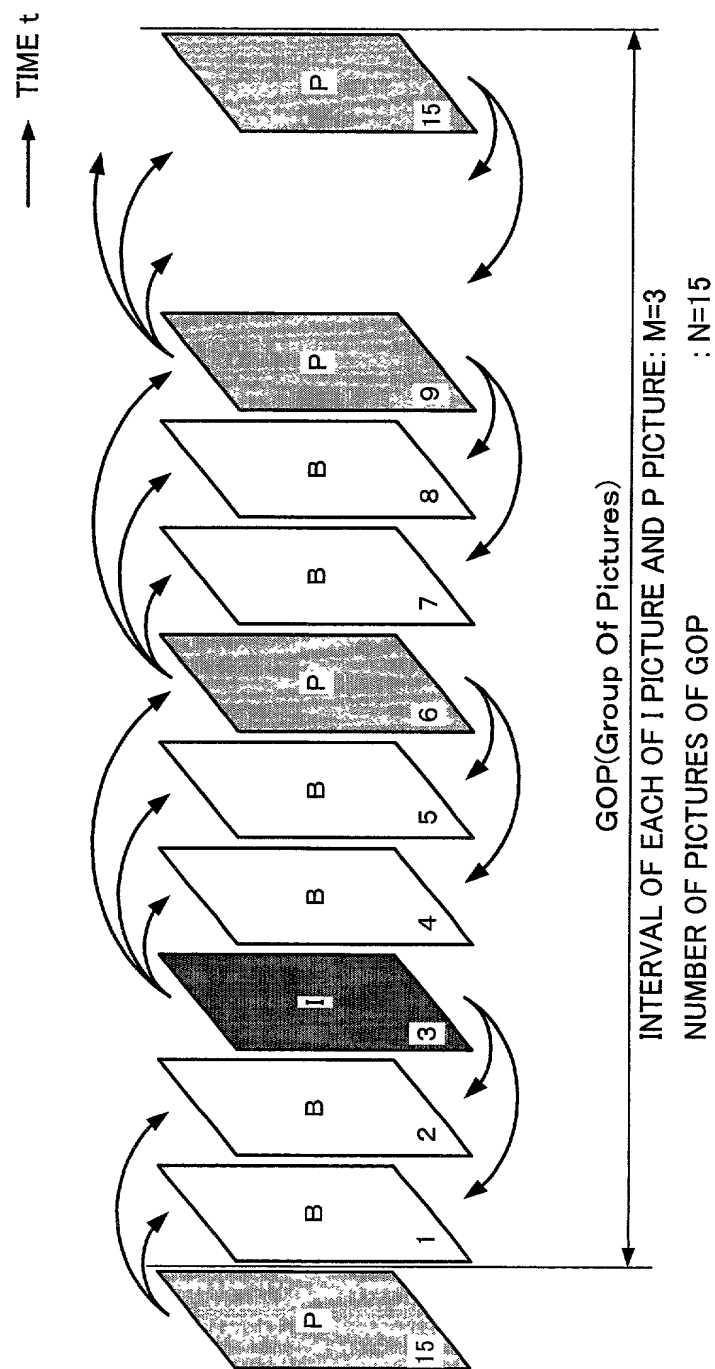


Fig. 6

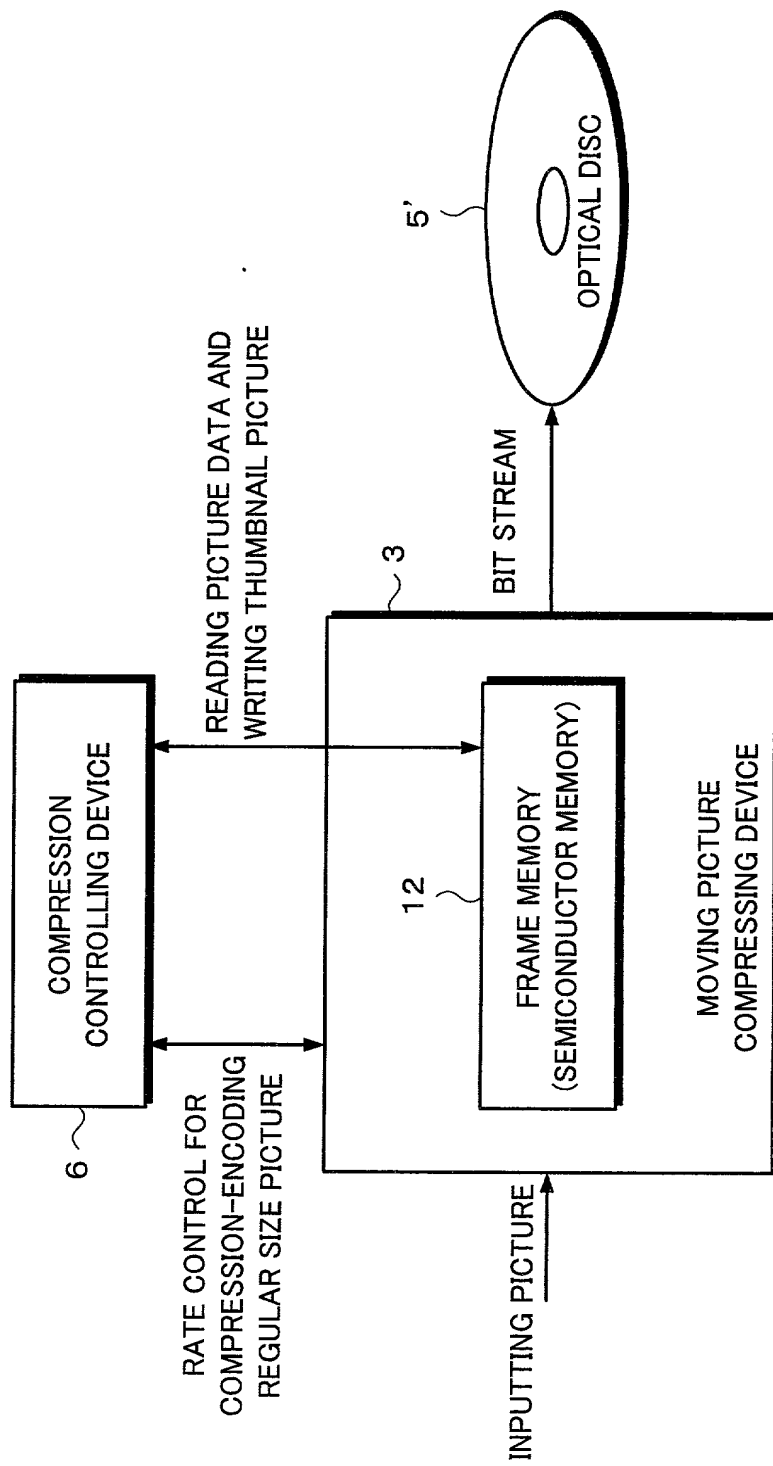


Figure 1 consists of 12 bar charts (a-l) showing the distribution of various parameters for the 1000 Genomes Project populations. The populations are: CEU (European), CHB (Chinese Han), CHS (Chinese Southern Han), JPT (Japanese), LWK (Luhya), MXL (Mexican), PUR (Purum), GBR (British), YRI (Yoruba), CHB+JPT (Chinese Han and Japanese), CHS+JPT (Chinese Southern Han and Japanese), and CEU+GBR (European and British). The parameters are: a) LD (r²), b) LD (r²), c) LD (r²), d) LD (r²), e) LD (r²), f) LD (r²), g) LD (r²), h) LD (r²), i) LD (r²), j) LD (r²), k) LD (r²), and l) LD (r²). Each chart shows the distribution of LD values for different population pairs.

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Fig. 8

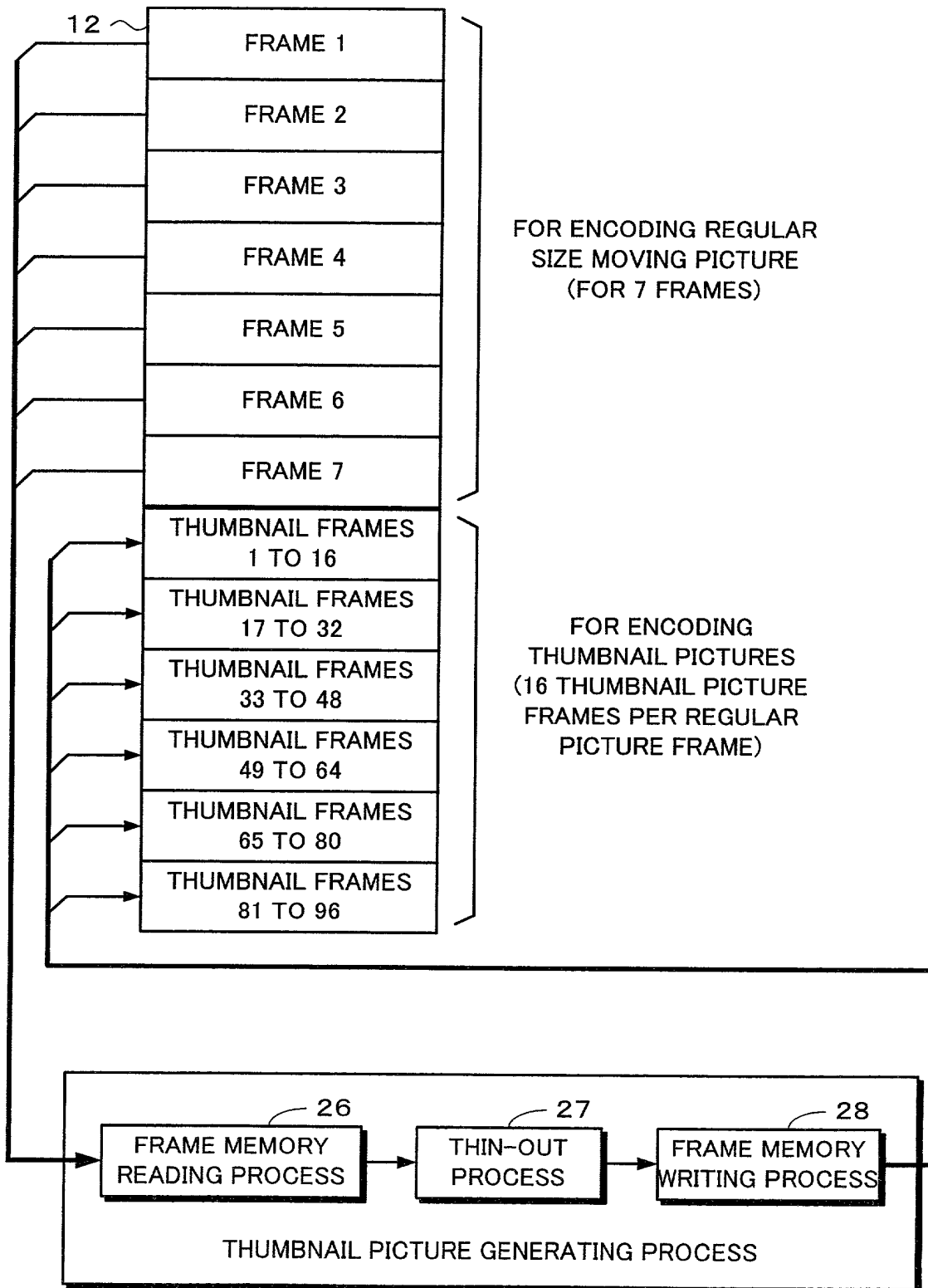


Fig. 10

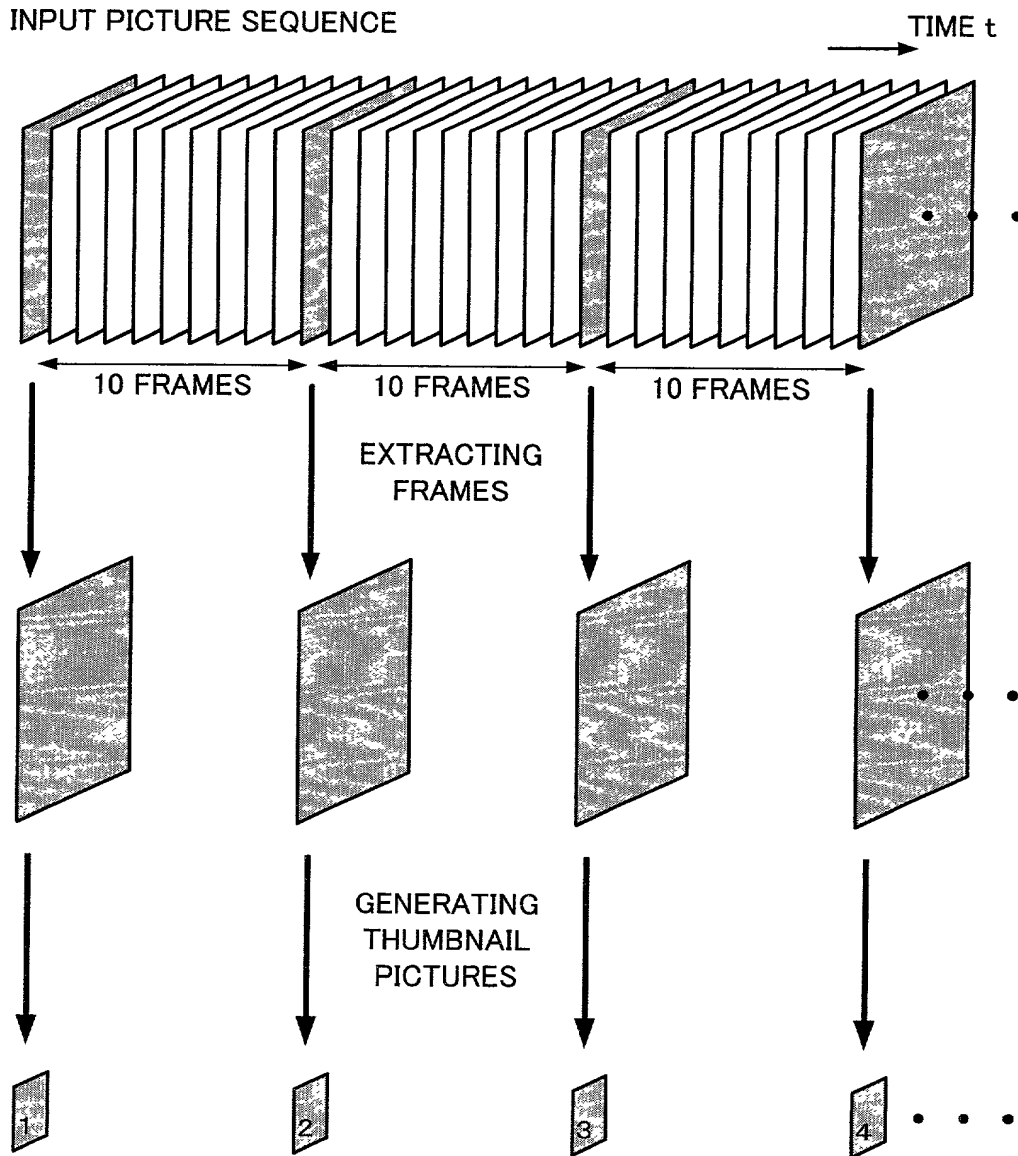


Fig. 11

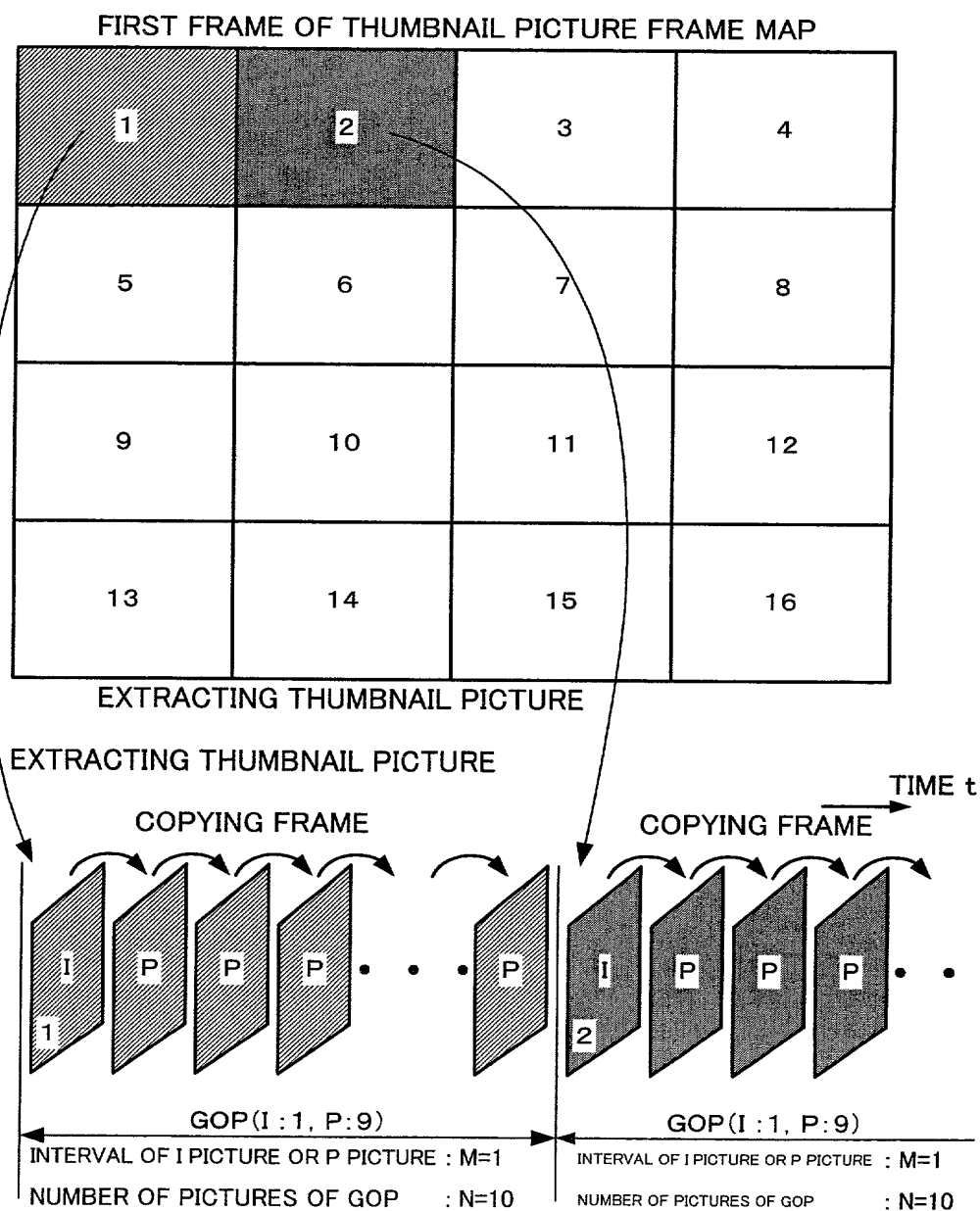
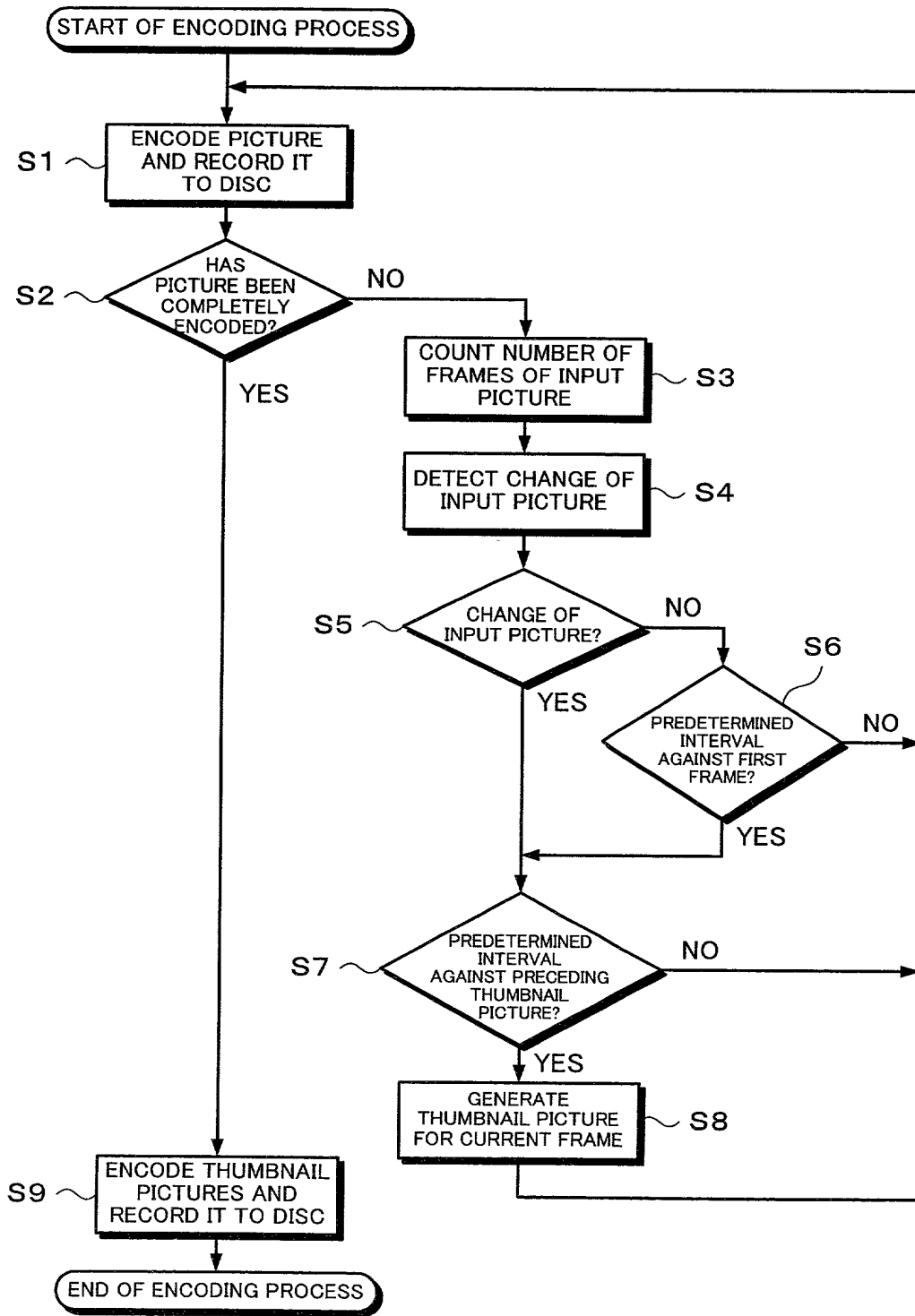


Fig. 12



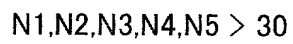


Fig. 14

FIRST FRAME OF THUMBNAIL PICTURE FRAME MAP

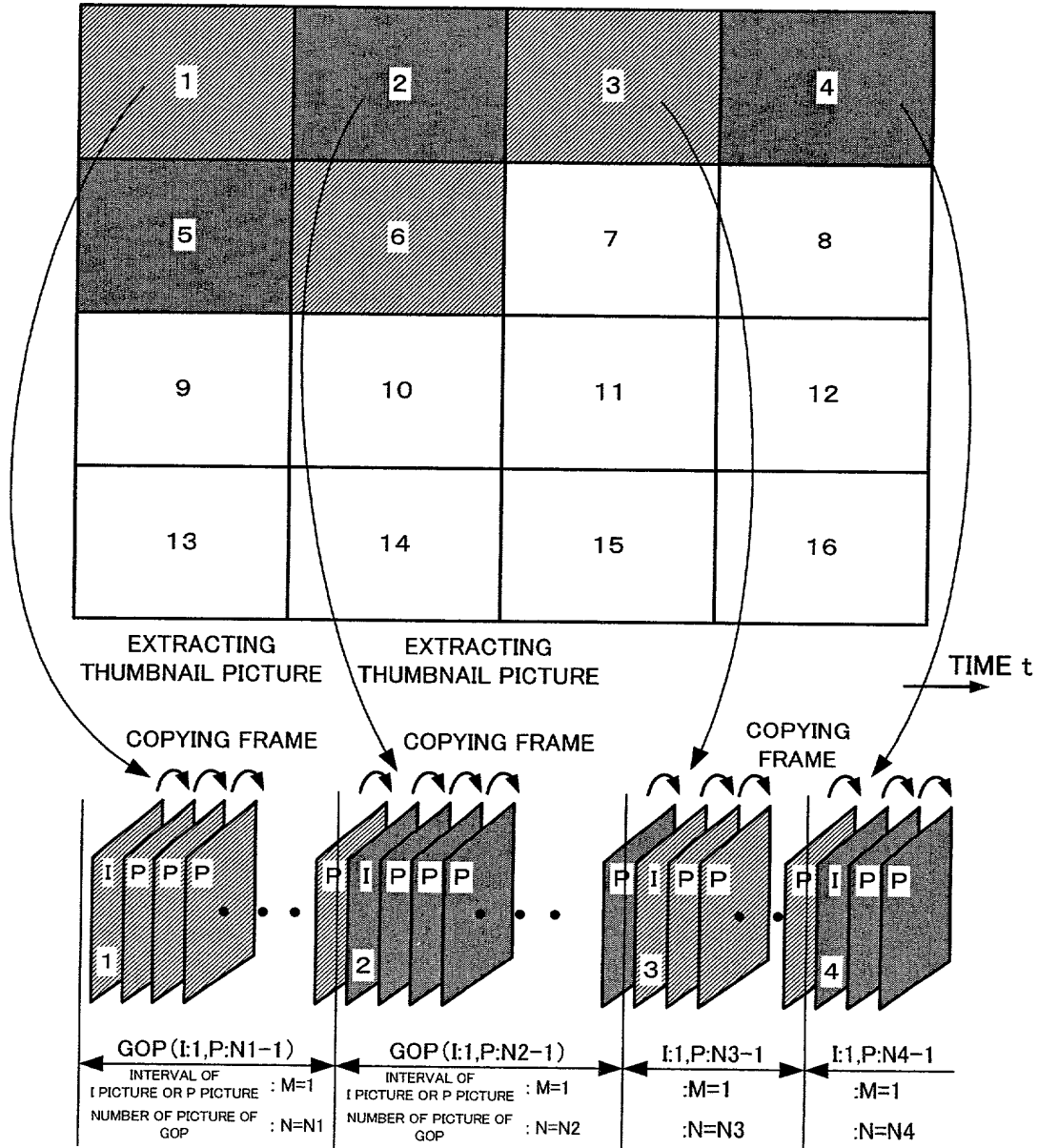


Fig. 16

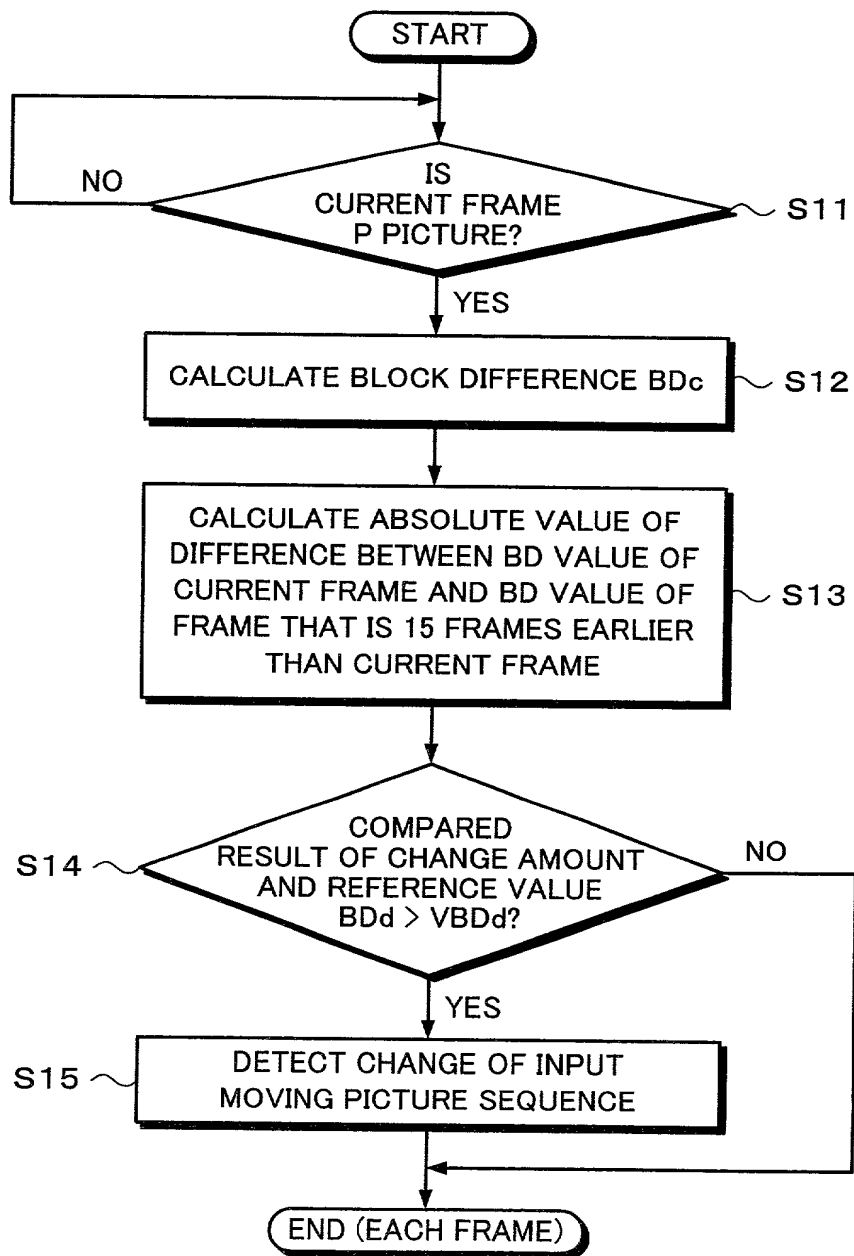


Fig. 17

